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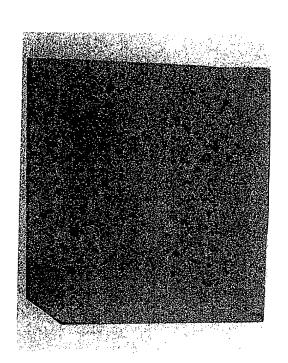
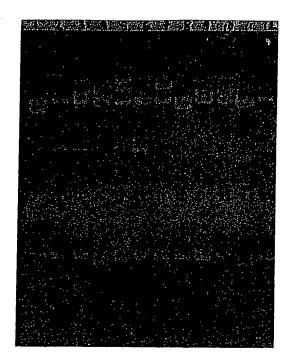


FIG.1



26 123

FIG.2

3/13

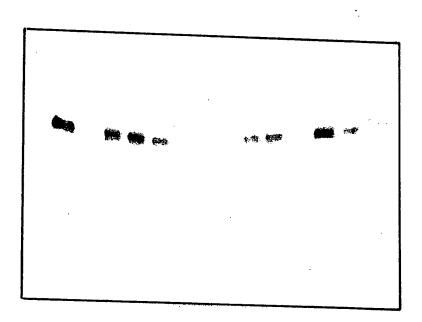


FIG.3



FIG.4

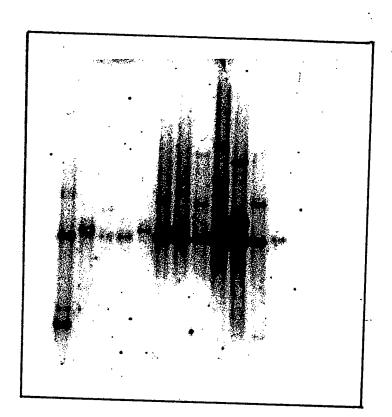


FIG.5



FIG.6

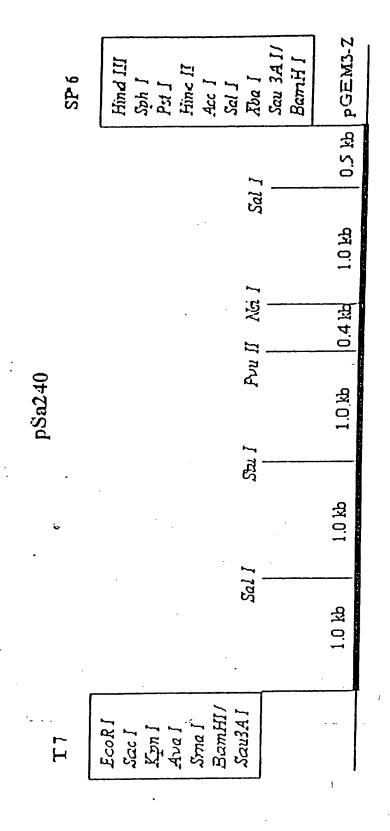


FIG.7

Fig. 8 Sa240 DNA sequence (4,826 bp)

GATCGGCGGCCGGTCGGCGGTGCTGGCCGCGGTGACCCTGGGGGGCGCTGGCCGCTCCGGCGGTGCTGCT GCGCCGGGGGCTGGCGGCCACCGCGGAGGCGCTGGCGGCGCTGGCCCTGGTGCTGACGCTGCTGGACGT GTACGCGGTCCACGCGGTGGCCGCGCCGGACACCGACGGACTCGGCTTCACGGCCCTCGCGTCGGCGGT GGTGCTGGCCCAGTGGCCGCTGCTGTTCTGGGCCTGGGCCGTGGGCGCACCGGCCGCCGGTGGTCGGGTGG GCGCTGCTGGCCACCGCGGTGCTGGACGGGGCGATCGCCCTGTGGGGCAAGGGCGCCGGGGTGCGGGTC ACGGCGTGCGTCGGTGAGCGCTGATGGGCTTCTCGGCCCTGATGGTGGGCCTGGCGCTGTCCCTGACGG CCGTCGGCGGCGTACCGGCGGCGCGCTCCCGGCGGGCTGGCGGGTGCTCGCGTACCTGCTGTGCGGTCT GGTGCTGTCGGACGTGTGGGCGGGGACGCCGGACGGCTTCCGGTCCGCGCTGGGGTCGACGCTGCCCTG OTCGGAGCTGGCCGCGGCCCCGGTGGTGCTCGCGCTGGTGGCCGGGCATGCTGGGGGGCGAGCGTAACCG GAGGTGGCCGTCGGTCCGGCTCCGGCGCCGTTGGCCGGTCCTTCTGGCTCGACGCCGGCCCCCCGGC TGGACGTGCCCCACGCGCTCGCGCTGGCCGGGGAGACGGCTCTGGTGGGCGTCCTGCTCGCCCTGGCGGT GGCGGTGAGCGCCGGGCTGCTGTCGCTGGCGTCCGAGGGGGCCTCGTACGCGGTGTTCGGCGCGCTCGC GGTGGTCTGGGGCACCGTGATCACGGGGTTGGCGGGCCGGTCCCTGGGGCTCGCCCCGCACGAGGCCGC CCCGCTGATGCTGCTGGTGCCGGCGCTGACGGTGCTGCTCGGGGCACGACTGCGGCGGAACCCGGTGGC CTTGCCCGTGGAGCTGACGGGAGCGCTGGGGCGCGCTCGTCGCCGTGGGGCTCGCGGTGTCCGACGCGCC GCCGGTGGCGGGCTACCTGGCGGCGACGCTGTTCGTGCTGGCCACGTGGGTGCGGCTGGCGGCCTCGGA GCGCCGGACCCGGAGGCCTCGTCGTGGACGGCGTACGGGCCGGGGGCTCGCGGCGACGCTGCTGCCCAG CCTGGCGGTCGCCTGGACCGACCGGACTGGCTCAGGCCGTTGCTGCTGGGGACGGCGGCGCTGGTGATC **ACCCTGCTCGGCGCGCCCACCGGCTCCAGGCGCTGCTGCTGCTCGGCGGGACGGTGCTGGCACTGGTCG** GGACGCGCTGGGGCGGATGCGGTGAGCCGTGCCCGGTCCGGGGGCGCGCAGGTCACGGCGTCCCCGGGC CCAGTGACCTCTTCGGTGTGAACGAAGCGCTCTCCCACTGAGCTAATCGCCCGGGCGCACCGCAAACATF ACCCCATGTCAGCGGTGCTCCCGGACCGTCCCCGGGCTACTCGCTGATCTTCCACGGCATGGTGAGCCCG AACTTCCAGACGTAGATCCCGGCCAGCACCGCCATGATCACGAGCCCGAGCGTGGTGAGGATGATGTTG CGCCGCCGGACCTTGGGATCGAGGGCCCGCTGCGCCGCTTCGGTGACCTTGCGCTTGGTCCAGCGCAGCA CCGGCAGCACCAGCATGAGCACACCCGCGATCACCACGCCGAGACCGATGAAGACACCGACCTGCC AGCTCAGGTGGAGCGCCTTGGACGCCTTGATGAAACCCGGCGCCCGCGAGCCCAGCGCGCGTTCCTCCCG GTCCGATTCCCCCGTGGCGGATACCGGGGACGCCTGCTCGGCGACCTTGCTCCGCTCGTCACTCTCCGCG TTCATGAAGCTCAACTTACCCGACCTGTCTCCGTCACTGGAATGGGCGCATAACTCAAAGTTACACGCCG CTGAGCGGGGGACCCGAAGCGTCACAAATGGGTCAGAGGGGTTTACAACGCCACCGTAGGTGGCATGTC GATTTCGCCGACGTGCGAATCCCCGAGCGCACACTGAGCGAAAAGGCCCTGGCGCTTATGAACACCACGG GTATGACACGGCCGATCCCTATGCCGTGCACGCCACCTTCCACACCGGAGCGGAGGAGACGGTCGAATG GGTATTCGCCCGCGACCTCCTTGCCGAGGGGCTGCACCGGCCCACCGGCACCGGAGACGTCCGCGTCTGG CCATCTCGTAGTCACGGTCAAGGCGTCGTATGCATCGCCCTGAGCTCCCCAGAGGGAGAAGCCCTGCTCG AAGCCCGGCGGGGCCCTGGAGTCGTTCCTGAAGAGGCCGACGCCGCGGTTCCGCCCGGCACCGAGC ATCGTCACTTCGATCTCGACACGGAGCTCTCCCACATCCTGGCCGAGAGCTGAGCCAGGCAGAGAGCCGC TCTACGCCGTCCGACTCGGGGCGACGGCGTCGTGCTGACAACCGCATAGGGCAGACACCGGCGCCGTCG TCGCGGAÀTCCACCGCGACGACGGCGCGCGCGCGTTCCCCGCCGCGCCGCGGAGGGGTCCGTTCCGCTC CCCGCCGCCGGAAGGCCAGGGAGCGAAGCGTGCTGATCCCTCACGACACCCGGATCGCCCTCGACGCG GGGCGGGCCGAGGACGGTCTCCCCGACATCGCCGCGCTGTACGCCTTCGCGGAGCGCCATCTCATCAGC GGGCTCGGCACCCTCGGCGAGAAGGACCTCGGCGCGCGTGCGCGACGTCCGGGCCCGCTTCGCCGAGGTC





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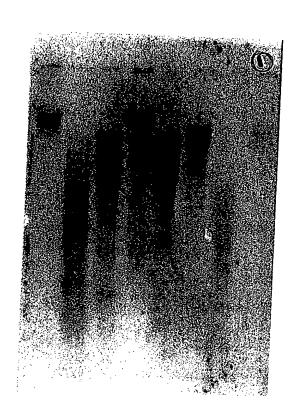
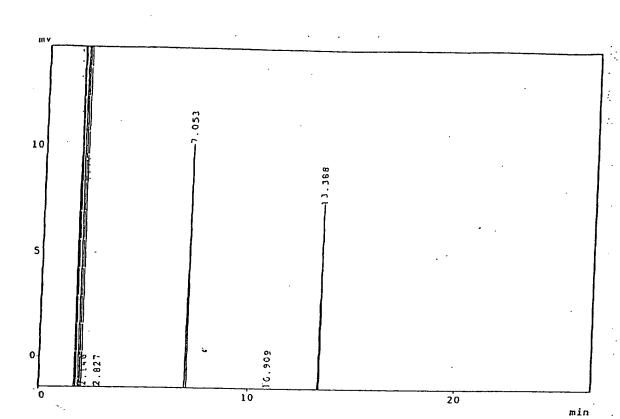


FIG.9



•••	Peak Report	t •••					
:010	TIME	AREA	HEIGHT	мк	IDNO	CONC	NAME
1	1.742	1090524	630697	V		16.30	
2	1.803	2759591	1055567	VE	41.2702 12.5883		
3	1.931	841734	440026	v			
4	1.989	1899444	1050304	SVE		28.406	
5	2.146	1222	1166	T		0.018	
6	2.027	1691	1181	v		0.025	
7	7.053	56479	14295			0.844	
8	10.909	1014	280			0.015	
9	13.388	34942	11651		·	0.522	
66		6686640	3205166			100.000	0

FIG.10

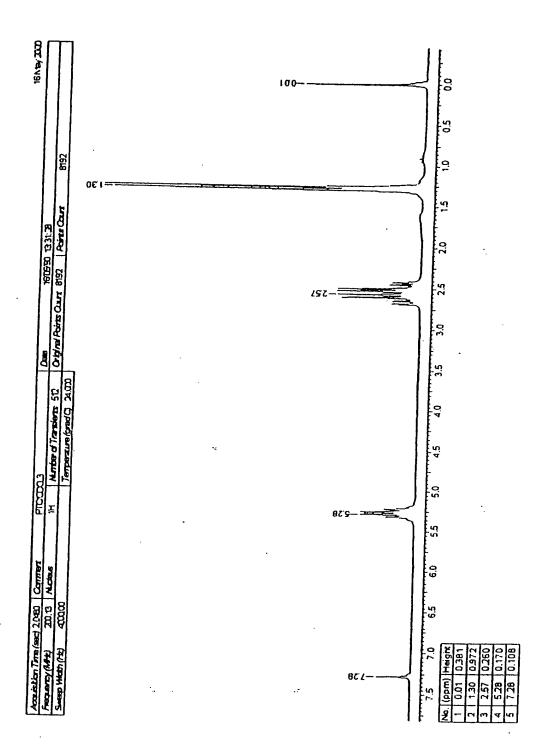
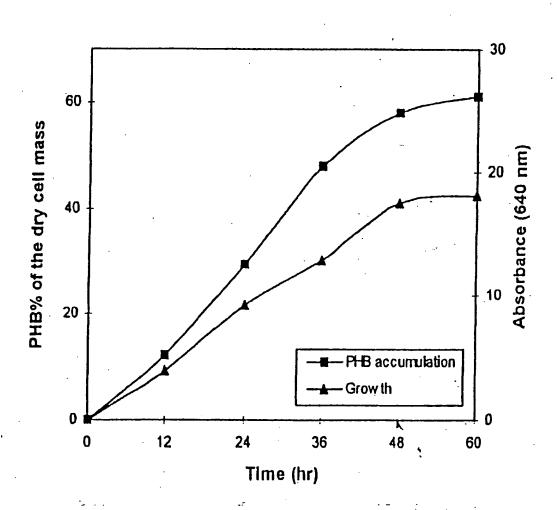


FIG.11



**FIG.12**